

Data transformation

Summarising and joining datasets

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05/10/2022

Let's recap

- tibbles
- Import and export tibbles
- filter data by row
- select columns (variables)
- pivot longer (columns to rows)
- pivot wider (rows to columns)

summarise()

Calculate the average age of sinai covid patients

```
# load libraries
library(tidyverse)
library(janitor)

# Import data
sinai_covid <- read_csv("sinai_covid.csv")

# clean names
sinai_covid <- sinai_covid %>%
    clean_names()
```

Even better, assign a name to the result

```
sinai_covid %>%
  summarise(mean_age = mean(age))

## # A tibble: 1 × 1
## mean_age
## <dbl>
## 1 61.3
```

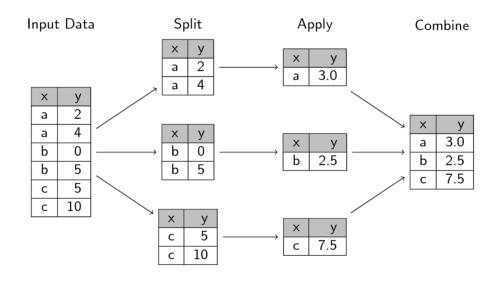
Calculate more than one summarise

Your turn!

• Use summarise to calculate the median, max and min values of age

How does summarise works?

Split-Apply-Combine



group_by()

Get the mean age and sd of sinai covid patients by sex

Your turn!

• Calculate the median age, and average bmi of sinai covid patients by facility

We can use multiple variables to group by

sinai covid %>%

```
group by (facility, sex) %>%
  summarise(mean age = mean(age),
            sd age = sd(age)) %>%
  head(4)
## `summarise()` has grouped output by 'facility'. You
## `.groups` argument.
## # A tibble: 4 \times 4
## # Groups: facility [2]
## facility
                              sex mean age sd ag
## <chr>
                              <chr> <dbl> <dbl
## 1 MOUNT SINAI BI BROOKLYN FEMALE 63.8 17.
## 2 MOUNT SINAI BI BROOKLYN MALE 64.6 14.
## 3 MOUNT SINAI QUEENS HOSPITAL FEMALE 69.2 14.
## 4 MOUNT SINAI QUEENS HOSPITAL MALE 62.8 14.
```

Let's practice!

• Find the mean, median and sd of the systolic_bp by smoking status and sex

count()

For qualitative variables

Excercise

• How many patients have chronic kidney disease?

group and count

```
sinai covid %>%
  group by (facility) %>%
  count (smoking status, diabetes)
## # A tibble: 27 \times 4
## # Groups: facility [5]
##
     facility
                                 smoking status diabe
##
  <chr>
                                 <chr>
                                                   < d
##
  1 MOUNT SINAI BI BROOKLYN
                                 NEVER
##
  2 MOUNT SINAI BI BROOKLYN NEVER
##
  3 MOUNT SINAI BI BROOKLYN QUIT
##
  4 MOUNT SINAI BI BROOKLYN QUIT
##
  5 MOUNT SINAI BI BROOKLYN
                             YES
##
  6 MOUNT SINAI QUEENS HOSPITAL NEVER
##
  7 MOUNT SINAI QUEENS HOSPITAL NEVER
##
  8 MOUNT SINAI QUEENS HOSPITAL QUIT
##
  9 MOUNT SINAI QUEENS HOSPITAL QUIT
##
  10 MOUNT SINAI QUEENS HOSPITAL YES
  # ... with 17 more rows
```

```
sinai_covid %>%
  group_by(facility) %>%
  filter(diabetes == 1) %>%
  count(smoking_status, diabetes)
```

```
## # A tibble: 13 \times 4
## # Groups: facility [5]
##
      facility
                                   smoking status diabe
##
     <chr>
                                   <chr>
                                                     < d
## 1 MOUNT SINAI BI BROOKLYN
                                  NEVER
## 2 MOUNT SINAI BI BROOKLYN
                                  QUIT
## 3 MOUNT SINAI BI BROOKLYN
                                  YES
## 4 MOUNT SINAI QUEENS HOSPITAL NEVER
## 5 MOUNT SINAI QUEENS HOSPITAL QUIT
##
  6 MOUNT SINAI ST. LUKE'S
                                  NEVER
##
  7 MOUNT SINAI ST. LUKE'S
                                  QUIT
##
  8 MOUNT SINAI WEST
                                  NEVER
##
  9 MOUNT SINAI WEST
                                  QUIT
##
  10 MOUNT SINAI WEST
                                  YES
##
  11 THE MOUNT SINAI HOSPITAL
                                  NEVER
##
   12 THE MOUNT SINAI HOSPITAL
                                   QUIT
```

join()

• Create tibble 1

```
tibble1 <- sinai_covid %>%
  filter(smoking_status %in% c("YES", "NEVER'
  count(smoking_status, name = "total")

tibble1
```

• Create tibble 2

- Join tibbles
- Keep rows from left tibble

```
left_join(tibble1, tibble2)

## Joining, by = "smoking_status"

## # A tibble: 2 × 4

## smoking_status total n_diabetes n_obesity
## <chr> <int> <dbl> <dbl> <dbl>
## 1 NEVER 341 NA NA
## 2 YES 29 6 3
```

• Keep rows from right tibble

• Keep rows in common

```
inner_join(tibble1, tibble2)

## Joining, by = "smoking_status"

## # A tibble: 1 × 4

## smoking_status total n_diabetes n_obesity
## <chr> <int> <dbl> <dbl> <dbl>
## 1 YES
29
6
3
```

Keep all rows

```
full join(tibble1, tibble2)
## Joining, by = "smoking status"
## # A tibble: 3 \times 4
## smoking status total n diabetes n obesity
## <chr>
            <int> <dbl> <dbl>
## 1 NEVER
                   341
                              NA
                                      NA
                                     3
## 2 YES
                   29
## 3 QUIT
                             43
                                       18
                    NA
```

Let's practice

- Count the number of patients by ethnicity, store the result in tibble 1
- Calculate the mean systolic_bp and mean diastolic_bp by ethnicity, store the result in tibble 2
- Join the tibbles

Thanks!



Ilustration by Allison Horst